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Sequence Listing was accepted.

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Reviewer: Anne Corrigan

Timestamp: [year=2009; month=10; day=20; hr=13; min=1; sec=48; ms=890; ]

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Application No: 09996069 Version No: 3.0

Input Set:

Output Set:

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Finished: 2009-10-01 19:46:33.984  
Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 9 ms  
Total Warnings: 12  
Total Errors: 0  
No. of SeqIDs Defined: 12  
Actual SeqID Count: 12

Error code	Error Description
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W 213	Artificial or Unknown found in <213> in SEQ ID (10)
W 213	Artificial or Unknown found in <213> in SEQ ID (11)
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# SEQUENCE LISTING

<110> MINERVA BIOTECHNOLOGIES CORPORATION  
Bamdad, Cynthia

<120> DIAGNOSTIC TUMOR MARKERS, DRUG SCREENING FOR TUMORIGENESIS  
INHIBITION, AND COMPOSITIONS AND METHODS FOR TREATMENT OF CANCER

<130> M1015-70071US

<140> 09996069

<141> 2001-11-27

<160> 12

<170> PatentIn version 3.5

<210> 1

<211> 39

<212> PRT

<213> Artificial Sequence

<220>

<223> Histidine-Tagged Truncated Receptor (His-TR)

<400> 1

Gly	Thr	Ile	Asn	Val	His	Asp	Val	Glu	Thr	Gln	Phe	Asn	Gln	Tyr	Lys
1				5					10					15	

Thr	Glu	Ala	Ala	Ser	Pro	Tyr	Asn	Leu	Thr	Ile	Ser	Asp	Val	Ser	Val
			20					25					30		

Ser	His	His	His	His	His	His
						35

<210> 2

<211> 51

<212> PRT

<213> Artificial Sequence

<220>

<223> His-PSMGFR peptides

<400> 2

Gly	Thr	Ile	Asn	Val	His	Asp	Val	Glu	Thr	Gln	Phe	Asn	Gln	Tyr	Lys
1				5					10					15	

Thr	Glu	Ala	Ala	Ser	Pro	Tyr	Asn	Leu	Thr	Ile	Ser	Asp	Val	Ser	Val
				20				25					30		

Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala His His His  
35 40 45

His His His  
50

<210> 3

<211> 54

<212> PRT

<213> Artificial Sequence

<220>

<223> "Extended Sequence of the MUC1 Growth Factor Receptor" (ESMGFR)

<400> 3

Val Gln Leu Thr Leu Ala Phe Arg Glu Gly Thr Ile Asn Val His Asp  
1 5 10 15

Val Glu Thr Gln Phe Asn Gln Tyr Lys Thr Glu Ala Ala Ser Pro Tyr  
20 25 30

Asn Leu Thr Ile Ser Asp Val Ser Val Ser Asp Val Pro Phe Pro Phe  
35 40 45

His His His His His His  
50

<210> 4

<211> 31

<212> PRT

<213> Artificial Sequence

<220>

<223> Histidine-Tagged Primary Sequence of the Interchain binding  
Region (His-PSIBR)

<400> 4

His His His His His His Gly Phe Leu Gly Leu Ser Asn Ile Lys Phe  
1 5 10 15

Arg Pro Gly Ser Val Val Val Gln Leu Thr Leu Ala Phe Arg Glu  
20 25 30

<210> 5

<211> 46

<212> PRT

<213> Artificial Sequence

<220>

<223> Histidine-Tagged Repeat Motif 2 (His-RM2)

<400> 5

Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly  
1 5 10 15

Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro  
20 25 30

Pro Ala His Gly Val Thr Ser Ala His His His His His His  
35 40 45

<210> 6

<211> 33

<212> PRT

<213> Artificial Sequence

<220>

<223> Truncated Receptor (TR)

<400> 6

Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys  
1 5 10 15

Thr Glu Ala Ala Ser Pro Tyr Asn Leu Thr Ile Ser Asp Val Ser Val  
20 25 30

Ser

<210> 7

<211> 45

<212> PRT

<213> Artificial Sequence

<220>

<223> Primary Sequence of the MUC1 Growth Factor Receptor (PSMGFR)

<400> 7

Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys  
1 5 10 15

Thr Glu Ala Ala Ser Pro Tyr Asn Leu Thr Ile Ser Asp Val Ser Val  
20 25 30

Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala  
35 40 45

<210> 8

<211> 25

<212> PRT

<213> Artificial Sequence

<220>

<223> Primary Sequence of the Interchain Binding Region) (PSIBR)

<400> 8

Gly Phe Leu Gly Leu Ser Asn Ile Lys Phe Arg Pro Gly Ser Val Val  
1 5 10 15

Val Gln Leu Thr Leu Ala Phe Arg Glu  
20 25

<210> 9

<211> 40

<212> PRT

<213> Artificial Sequence

<220>

<223> Repeat Motif 2 (RM2)

<400> 9

Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly  
1 5 10 15

Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro  
20 25 30

Pro Ala His Gly Val Thr Ser Ala  
35 40

<210> 10

<211> 1255

<212> PRT

<213> Artificial Sequence

<220>

<223> Mucin 1 Precursor, Genbank Accession Number: P15941

<400> 10

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr

1	5	10	15
Val Leu Thr	Val Val Thr	Gly Ser Gly His Ala Ser Ser	Thr Pro Gly
	20	25	30
Gly Glu Lys	Glu Thr Ser Ala Thr	Gln Arg Ser Ser Val	Pro Ser Ser
	35	40	45
Thr Glu Lys	Asn Ala Val Ser Met Thr	Ser Ser Val Leu Ser	Ser His
	50	55	60
Ser Pro Gly	Ser Gly Ser Ser Thr Thr	Gln Gly Gln Asp Val Thr	Leu
65	70	75	80
Ala Pro Ala	Thr Glu Pro Ala Ser Gly	Ser Ala Ala Thr Trp Gly	Gln
	85	90	95
Asp Val Thr	Ser Val Pro Val Thr Arg	Pro Ala Leu Gly Ser Thr	Thr
	100	105	110
Pro Pro Ala	His Asp Val Thr Ser Ala	Pro Asp Asn Lys Pro Ala	Pro
	115	120	125
Gly Ser Thr	Ala Pro Pro Ala His Gly	Val Thr Ser Ala Pro Asp	Thr
	130	135	140
Arg Pro Ala	Pro Gly Ser Thr Ala Pro	Pro Ala His Gly Val Thr	Ser
145	150	155	160
Ala Pro Asp	Thr Arg Pro Ala Pro Gly	Ser Thr Ala Pro Pro Ala	His
	165	170	175
Gly Val Thr	Ser Ala Pro Asp Thr Arg	Pro Ala Pro Gly Ser Thr	Ala
	180	185	190
Pro Pro Ala	His Gly Val Thr Ser Ala	Pro Asp Thr Arg Pro Ala	Pro
	195	200	205
Gly Ser Thr	Ala Pro Pro Ala His Gly	Val Thr Ser Ala Pro Asp	Thr
	210	215	220
Arg Pro Ala	Pro Gly Ser Thr Ala Pro	Pro Ala His Gly Val Thr	Ser
225	230	235	240

Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His  
245 250 255

Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala  
260 265 270

Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro  
275 280 285

Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr  
290 295 300

Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser  
305 310 315 320

Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His  
325 330 335

Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala  
340 345 350

Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro  
355 360 365

Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr  
370 375 380

Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser  
385 390 395 400

Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His  
405 410 415

Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala  
420 425 430

Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro  
435 440 445

Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr  
450 455 460



Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser			
465	470	475	480
Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His			
	485	490	495
Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala			
	500	505	510
Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro			
	515	520	525
Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr			
	530	535	540
Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser			
545	550	555	560
Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His			
	565	570	575
Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala			
	580	585	590
Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro			
	595	600	605
Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr			
	610	615	620
Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser			
625	630	635	640
Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His			
	645	650	655
Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala			
	660	665	670
Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro			
	675	680	685

Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr  
690 695 700

Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser  
705 710 715 720

Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His  
725 730 735

Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala  
740 745 750

Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro  
755 760 765

Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr  
770 775 780

Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser  
785 790 795 800

Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His  
805 810 815

Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala  
820 825 830

Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro  
835 840 845

Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr  
850 855 860

Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser  
865 870 875 880

Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His  
885 890 895

Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala  
900 905 910

Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro

915

920

925

Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn  
 930 935 940

Arg Pro Ala Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser  
 945 950 955 960

Ala Ser Gly Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly  
 965 970 975

Thr Ser Ala Arg Ala Thr Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe  
 980 985 990

Ser Ile Pro Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His  
 995 1000 1005

Ser Thr Lys Thr Asp Ala Ser Ser Thr His His Ser Ser Val Pro  
 1010 1015 1020

Pro Leu Thr Ser Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr  
 1025 1030 1035

Gly Val Ser Phe Phe Phe Leu Ser Phe His Ile Ser Asn Leu Gln  
 1040 1045 1050

Phe Asn Ser Ser Leu Glu Asp Pro Ser Thr Asp Tyr Tyr Gln Glu  
 1055 1060 1065

Leu Gln Arg Asp Ile Ser Glu Met Phe Leu Gln Ile Tyr Lys Gln  
 1070 1075 1080

Gly Gly Phe Leu Gly Leu Ser Asn Ile Lys Phe Arg Pro Gly Ser  
 1085 1090 1095

Val Val Val Gln Leu Thr Leu Ala Phe Arg Glu Gly Thr Ile Asn  
 1100 1105 1110

Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys Thr Glu Ala  
 1115 1120 1125

Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val Ser Val Ser Asp  
 1130 1135 1140

Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala Gly Val Pro Gly  
1145 1150 1155

Trp Gly Ile Ala Leu Leu Val Leu Val Cys Val Leu Val Ala Leu  
1160 1165 1170

Ala Ile Val Tyr Leu Ile Ala Leu Ala Val Cys Gln Cys Arg Arg  
1175 1180 1185

Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr  
1190 1195 1200

His Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg Tyr  
1205 1210 1215

Val Pro Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Ser  
1220 1225 1230

Ala Gly Asn Gly Gly Ser Ser Leu Ser Tyr Thr Asn Pro Ala Val  
1235 1240 1245

Ala Ala Ala Ser Ala Asn Leu  
1250 1255

<210> 11

<211> 302

<212> PRT

<213> Artificial Sequence

<220>

<223> Proopiomelanocortin  
(adrenocorticotropin/beta-lipotropin/alpha-melanocyte  
stimulating hormone/beta-melanocyte stimulating  
hormone/beta-endorphin) [Homo sapiens]. Accession number:

<400> 11

Ala Ala Ala Lys Glu Gly Lys Lys Ser Arg Asp Arg Glu Arg Pro Pro  
1 5 10 15

Ser Val Pro Ala Leu Arg Glu Gln Pro Pro Glu Thr Glu Pro Gln Pro  
20 25 30

Ala Trp Lys Met Pro Arg Ser Cys Cys Ser Arg Ser Gly Ala Leu Leu

35

40

45

Leu Ala Leu Leu Leu Gln Ala Ser Met Glu Val Arg Gly Trp Cys Leu  
50 55 60

Glu Ser Ser Gln Cys Gln Asp Leu Thr Thr Glu Ser Asn Leu Leu Glu  
65 70 75 80

Cys Ile Arg Ala Cys Lys Pro Asp Leu Ser Ala Glu Thr Pro Met Phe  
85 90 95

Pro Gly Asn Gly Asp Glu Gln Pro Leu Thr Glu Asn Pro Arg Lys Tyr  
100 105 110

Val Met Gly His Phe Arg Trp Asp Arg Phe Gly Arg Arg Asn Ser Ser  
115 120 125

Ser Ser Gly Ser Ser Gly Ala Gly Gln Lys Arg Glu Asp Val Ser Ala  
130 135 140

Gly Glu Asp Cys Gly Pro Leu Pro Glu Gly Gly Pro Glu Pro Arg Ser  
145 150 155 160

Asp Gly Ala Lys Pro Gly Pro Arg Glu Gly Lys Arg Ser Tyr Ser Met  
165 170 175

Glu His Phe Arg Trp Gly Lys Pro Val Gly Lys Lys Arg Arg Pro Val  
180 185 190

Lys Val Tyr Pro Asn Gly Ala Glu Asp Glu Ser Ala Glu Ala Phe Pro  
195 200 205

Leu Glu Phe Ly